

SEQUENCE LISTING

<110> WOLFFE, Alan
URNOV, Fyodor
GUSCHIN, Dmitry
COLLINGWOOD, Trevor
LI, Xiao-Yong
JOHNSTONE, Brian

<120> DATABASES OF REGULATORY SEQUENCES; METHODS OF MAKING AND USING SAME

<130> 8325-0015

<140> 09/844,501
<141> 2001-04-27

<150> 60/200,590
<151> 2000-04-28

<150> 60/214,674
<151> 2000-06-27

<150> 60/228,556
<151> 2000-08-28

<160> 24

<170> PatentIn Ver. 2.0

<210> 1
<211> 6
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Kpn 1 target site

<400> 1
ggtacc

6

<210> 2
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: adapter oligonucleotide

<400> 2
gcggtgaccc gggagatctg aattc

25

<210> 3

<211> 11
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: adapter
oligonucleotide

<400> 3
ctagacttaa g 11

<210> 4
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Bax
gene-specific primer

<400> 4
gcccatcaact gagaaaatccc ttcc 24

<210> 5
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: adapter
oligonucleotide

<400> 5
gcggtgaccc gggagatctg aattctt 27

<210> 6
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: adapter
oligonucleotide

<400> 6
cgccactggg ccctctagac ttaag 25

<210> 7
<211> 60
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: adapter

oligonucleotide

<400> 7
tagaaggcac agtcgaggac ttatcctagc ctctgaatac tttcaacaag ttacaccctt 60

<210> 8
<211> 66
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: adapter
oligonucleotide

<400> 8
aaaaaaaaatc ttccgtgtca gctcctgaat aggatcgagg acttatgaaa gttgttcaat 60
gtggga 66

<210> 9
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence:
adapter-specific primer

<400> 9
aggcacagtc gaggacttat ccta 24

<210> 10
<211> 122
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: insert
sequence

<400> 10
ccggcctcgg tgttttcggc ttttcctgg cccccggccc gccaggccgg gccctctgct 60
gcccgcgtaa tgggagggggg ggccgggtca cgtggcgggg ggagggggagg gccgtcgcgta 120
tc 122

<210> 11
<211> 249
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: insert
sequence

<400> 11

ccgggcgcca aggaaagccg ggcgctgcc cctgctggcc aggttcgggc gcggcgccgc 60
ggaggggcct cccctctcg gagagaattg aagggggtcc ggtgtggagc cccggctggc 120
tccgggctgg ggctgaccgg ctctgtgacc ttgggcaggt cactgcatct ctccaagcct 180
cagtttgcac gtctgtcaaa tagagggca ttctctact ttgcagggtc cctggaaata 240
agttagatc 249

<210> 12
<211> 1042
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: accessible
region sequence

<400> 12
gatcgaggt cgagaccagc ccggccaact ggtgaaaccc tgcgttact aaaaaaatac 60
aaaaggagtt cgagaccagc ccggccaact ggtgaaaccc tgcgttact aaaaaaatac 120
aaaaatttagc tgggtgttgtt ggtgcacgcc tgcgttact aaaaaaatac 180
aggaatttagc tgggtgttgtt ggtgcacgcc tgcgttact aaaaaaatac 240
aggagaatcg cttgaaccca ggaggggagg cagagggtgc agtgagccga gatggcgcca 300
ctgtgaatcg cttgaaccca ggaggggagg cagagggtgc agtgagccga gatggcgcca 360
ctgtactccg gcctgggcaa gagcaagact ccaaccaaaa aaaaaaaaaa aaagaactag 420
cagtaactccg gcctgggcaa gagcaagact ccaaccaaaa aaaaaaaaaa aaagaactag 480
cagtgcccag ggctgtacac caggtgccag tactggcagc aattcttcca gttattgtga 540
tagagcccag ggctgtacac caggtgccag tactggcagc aattcttcca gttattgtga 600
tagattctca tgacgctaaa atacccactt tgttatttaa cccttgctaa tccacaatga 660
gttgttctca tgacgctaaa atacccactt tgttatttaa cccttgctaa tccacaatga 720
gttgcaggt accagaatcc ttgttacta accagaccag gctgttcatt cttgaacagc 780
attgccaggt accagaatcc ttgttacta accagaccag gctgttcatt cttgaacagc 840
attgggcattc actttgttt aataattctt gtatgagaag agcaactctt tccttctgat 900
agcaggcatc actttgttt aataattctt gtatgagaag agcaactctt tccttctgat 960
agcaatgtgg ctccaaactac tggctgatgt gagacggtac cggatgtggc tccaaactact 1020
ggctgatgtg agacggtacc gg 1042

<210> 13
<211> 12
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: adapter
oligonucleotide containing a Sau 3AI-compatible
end

<400> 13
gatcgaattc ag 12

<210> 14
<211> 8
<212> DNA
<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: adapter oligonucleotide containing a Sau 3AI-compatible end	
<400> 14 cttaagtc	8
<210> 15 <211> 20 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: p16 forward primer	
<400> 15 aatagcacct cctccgagca	20
<210> 16 <211> 21 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: p16 reverse primer	
<400> 16 ccctgtccct caaatcctct g	21
<210> 17 <211> 23 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: p16 probe	
<400> 17 acagcgcccc cttgcctggaa aag	23
<210> 18 <211> 19 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Control forward primer	
<400> 18 gccccagagg gaaacaccaa	19

<210> 19		
<211> 17		
<212> DNA		
<213> Artificial Sequence		
<220>		
<223> Description of Artificial Sequence: Control		
reverse primer		
<400> 19		
ccccccaccccccataaggc		17
<210> 20		
<211> 24		
<212> DNA		
<213> Artificial Sequence		
<220>		
<223> Description of Artificial Sequence: Control probe		
<400> 20		
cctccatgggt ggtaccacagc aagg		24
<210> 21		
<211> 48		
<212> DNA		
<213> Artificial Sequence		
<220>		
<223> Description of Artificial Sequence: EPAS		
amplifier primer		
<400> 21		
ggatccggcc accgcggccg cacgccccat agccctgaag actattac		48
<210> 22		
<211> 44		
<212> DNA		
<213> Artificial Sequence		
<220>		
<223> Description of Artificial Sequence: EPAS		
amplifier primer		
<400> 22		
atgaatttcgc ggccggccca ctgggtattt gatctggccc ccat		44
<210> 23		
<211> 109		
<212> DNA		
<213> Artificial Sequence		
<220>		
<223> Description of Artificial Sequence: human VEGF		

accessible region

<400> 23
atcagagaca ggctctgtct gccagctgtc tctccctcag ggctctgcc aactccacag 60
tgcatacgtg ggcttccaca ggtcgctcc ctccggcac tgactaact 109

<210> 24
<211> 134
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: human VEGF
accessible region

<400> 24
catctgggt tggggggca gcaggaacaa gggccttgt ctgcccagct gcctccccct 60
ttgggttttg ccagactcca cagtgcatac gtgggctcca acaggtcctc ttccctccca 120
gtcactgact aacc 134